



ENTERED

1600

RAW SEQUENCE LISTING

DATE: 08/11/2003

PATENT APPLICATION: US/09/776,705B

TIME: 12:31:16

Input Set : A:\1010 SEQ LISTING.TXT

Output Set: N:\CRF4\08112003\I776705B.raw

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4 <110> APPLICANT: Karl GUESLER et al.
6 <120> TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
7 NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
8 AND USES THEREOF
10 <130> FILE REFERENCE: CLO11010
11 <140> CURRENT APPLICATION NUMBER: 09/776,705B
13 <141> CURRENT FILING DATE: 2001-02-06
15 <150> PRIOR APPLICATION NUMBER: 60/251,836
16 <151> PRIOR FILING DATE: 2000-12-03
18 <160> NUMBER OF SEQ ID NOS: 73
20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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24 <212> TYPE: DNA
25 <213> ORGANISM: Homo Sapiens
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30 cagaaggggc ttctgtgttg acaagaaagc gctgacagct caaatggatc ccattggaac 160
31 gagaaatgtc aacatcgaac cagatgatga gagcagcagt ggagaaagtg ctccagatag 210
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62 <312> TYPE: PRT

63 <313> ORGANISM: Homo Sapiens

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70 35 40 45
71 Ser Gln Lys Phe Leu Thr Asn Gly Phe Leu Gly Lys Lys Lys Leu Ala
72 50 55 60
73 Asp Tyr Ala Asp Glu His His Pro Gly Thr Thr Ser Phe Gly Met Ser
74 65 70 75 80
75 Ser Phe Asn Leu Ser Asn Ala Ile Met Gly Ser Gly Ile Leu Gly Leu
76 85 90 95
77 Ser Tyr Ala Met Ala Tyr Thr Gly Val Ile Leu Phe Ile Ile Met Leu
78 100 105 110
79 Leu Ala Val Ala Ile Leu Ser Leu Tyr Ser Val His Leu Leu Leu Lys
80 115 120 125
81 Thr Ala Lys Glu Gly Gly Ser Leu Ile Tyr Glu Lys Leu Gly Glu Lys
82 130 135 140
83 Ala Phe Gly Trp Pro Gly Lys Ile Gly Ala Phe Val Ser Ile Thr Met
84 145 150 155 160
85 Gln Asn Ile Gly Ala Met Ser Ser Tyr Leu Phe Ile Ile Lys Tyr Glu
86 165 170 175
87 Leu Pro Glu Val Ile Arg Ala Phe Met Gly Leu Glu Glu Asn Thr Gly
88 180 185 190
89 Glu Trp Tyr Leu Asn Gly Asn Tyr Leu Ile Ile Phe Val Ser Val Gly
90 195 200 205
91 Ile Ile Leu Pro Leu Ser Leu Leu Lys Asn Leu Gly Tyr Leu Gly Tyr
92 210 215 220
93 Thr Ser Gly Phe Ser Leu Thr Cys Met Val Phe Phe Val Ser Val Val
94 225 230 235 240
95 Ile Tyr Lys Lys Phe Gln Ile Pro Cys Pro Leu Pro Val Leu Asp His
96 245 250 255
97 Ser Val Gly Asn Leu Ser Phe Asn Asn Thr Leu Pro Met His Val Val
98 260 265 270
99 Met Leu Pro Asn Asn Ser Glu Ser Ser Asp Val Asn Phe Met Met Asp
100 275 280 285
101 Tyr Thr His Arg Asn Pro Ala Gly Leu Asp Glu Asn Gln Ala Lys Gly
102 290 295 300
103 Ser Leu His Asp Ser Gly Val Glu Tyr Glu Ala His Ser Asp Asp Lys
104 305 310 315 320

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110 Tyr Ser Glu Leu Lys Asp Arg Ser Arg Arg Lys Met Gln Thr Val Ser
111           355           360           365
112 Asn Ile Ser Ile Thr Gly Met Leu Val Met Tyr Leu Leu Ala Ala Leu
113           370           375           380
114 Phe Gly Tyr Leu Thr Phe Tyr Gly Glu Val Glu Asp Glu Leu Leu His
115           385           390           395           400
116 Ala Tyr Ser Lys Val Tyr Thr Leu Asp Ile Pro Leu Leu Met Val Arg
117           405           410           415
118 Leu Ala Val Leu Val Ala Val Thr Glu Thr Val Pro Ile Val Leu Phe
119           420           425           430
120 Pro Ile Arg Thr Ser Val Ile Thr Leu Leu Phe Pro Lys Arg Pro Phe
121           435           440           445
122 Ser Trp Ile Arg His Phe Leu Ile Ala Ala Val Leu Ile Ala Leu Asn
123           450           455           460
124 Asn Val Leu Val Ile Leu Val Pro Thr Ile Lys Tyr Ile Phe Gly Phe
125           465           470           475           480
126 Ile Gly Ala Ser Ser Ala Thr Met Leu Ile Phe Ile Leu Pro Ala Val
127           485           490           495
128 Phe Tyr Leu Lys Leu Val Lys Lys Glu Thr Phe Arg Ser Pro Gln Lys
129           500           505           510
130 Val Gly Ala Leu Ile Phe Leu Val Val Gly Ile Phe Phe Met Ile Gly
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134 Lys His His
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136 <10> SEQ ID NO: 3

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138 <12> TYPE: DNA

139 <13> ORGANISM: Homo Sapiens

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147 ttaaaaagcga ctacacatcta gtcacgtcgt gatgaaagga taaggataaa aattctgaaa 240
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149 agctagaaga gagaagtttc ttcaatatcc tgaaggaaaa tgcttctgaa totagaattc 360
150 aaacaattaa caaagtttga aggcacaaata aagaattttc caacatgaag caactcagaa 420
151 attctattta cagacatagg ctcatctgtt gaaaaaagtt attcaaggca ttatttttagc 480
152 ataattgcana ataaactgaa gaaagaagat agaattgcgt tcaagaaact agcagctgag 540
153 caagactcag aggttgagg aggaagccat tcagaatgag aaagagcata gaaaatttgc 600
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155 ttgtcttttag ggcatactat ttatacaactg ataatactgt aattgctgct tattggtttt 720
156 cratgttttag aaacaacctt caggcaagtt atgacacttg ttccacagaa caagatgaaa 780
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160 ttttaaglaac aagctaaaaa cagaaacogt caatagtggg tggcgctggg aagtggagct 820
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VERIFICATION SUMMARY

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STATISTICS SUMMARY

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Input Set : A:\1010 SEQ LISTING.TXT

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Application Serial Number: US/09/776,705B

Alpha or Numeric or Xml: Numeric

Application Class:

Application File Date: 02-06-2001

Art Unit: 1600

Software Application: FastSEQ

Total Number of Sequences: 78

Total Nucleotides: 67094

Total Amino Acids: 647

Number of Errors: 0

Number of Warnings: 0

Number of Corrections: 0

MESSAGE SUMMARY